

Promotor region



-1237 T>C

+2848 G>A

Wild type-1237/T GGGGGAGTTTT**T**CAGGCAGAGG

Mutant -1237/C GGGGGAGTTTT**C**CAGGCAGAGG

NF-kB consensus sequence GGGRNNYY**C**C

RelA consensus sequence GGRNWTT**C**C

STAT3 consensus sequence TT**C**CDGGAAR